

**Fig. 1**



**Fig. 3**

Name	DNAzyme Sequenz
hgd1	5'-TCGGTCAGAggctagctacaacgaTGCGTTGCT-3'
hgd2	5'-GGCGTACGAggctagctacaacgaCTGCTCGGT-3'
hgd3	5'-GGCGGCGTAGgctagctacaacgaGACCTGCTC-3'
hgd4	5'-CTCGGGTCAGgctagctacaacgaCTGGGTAGC-3'
hgd5	5'-TCCTCTGCAGgctagctacaacgaCGGGGTCCT-3'
hgd6	5'-ACTCTGCAAggctagctacaacgaTCTGCGAGC-3'
hgd7	5'-GGGCGACGAggctagctacaacgaTCTGCAATT-3'
hgd8	5'-AAGGGGCGAggctagctacaacgaGACTCTGCA-3'
hgd9	5'-AAAACGGGAggctagctacaacgaCAGGTTGTA-3'
hgd10	5'-AGAATAAAAggctagctacaacgaGGGACCAGG-3'
hgd11	5'-ATGGCAGAAggctagctacaacgaAAAACGGGA-3'
hgd12	5'-AACTGGGTAggctagctacaacgaGGCAGAATA-3'
hgd13	5'-ATCCAAAAAggctagctacaacgaTGGGTATGG-3'
hgd14	5'-AGGGGAAGAggctagctacaacgaAAAAATCCA-3'
hgd15	5'-TTTTAAAAAggctagctacaacgaTATCTTGGA-3'
hgd16	5'-GTGGGGGGAggctagctacaacgaGGGAAGGCT-3'
hgd17	5'-GTTGAATGAggctagctacaacgaTTGCTTTCG-3'
hgd18	5'-GTCGTTGAaggctagctacaacgaGATTTGCTT-3'
hgd19	5'-GGCCCCGAaggctagctacaacgaCCGCGCGCG-3'
hgd20	5'-TCACCTCCAaggctagctacaacgaGGCCTCGGC-3'
hgd21	5'-CCGCCGTCaggctagctacaacgaCTCCATGGC-3'
hgd22	5'-GGTGGCTCaggctagctacaacgaCCAGCGCGG-3'
hgd23	5'-CGTTGAGCaggctagctacaacgaGGCGGGGTG-3'
hgd24	5'-CCGCGTCCAaggctagctacaacgaGTAGGAGTG-3'
hgd25	5'-CAGCGGGTAggctagctacaacgaTGCGCCGCG-3'
hgd26	5'-GCACATCCAaggctagctacaacgaCTCCTCCGG-3'
hgd27	5'-AAAAGCACaggctagctacaacgaCCACCTCCT-3'
hgd28	5'-TAAAAAGCaggctagctacaacgaATCCACCTC-3'
hgd29	5'-GACCGTCGaggctagctacaacgaGTTAAAAAG-3'
hgd30	5'-TTGCCTTGaggctagctacaacgaCGTCGATGT-3'
hgd31	5'-AGGGCGGGAggctagctacaacgaGTGGTTGCC-3'
hgd32	5'-TGGCCCTGAggctagctacaacgaCGAGTTTCC-3'
hgd33	5'-ACCTCTGCAggctagctacaacgaCGTGGCCCT-3'
hgd34	5'-CGGAGGGTAggctagctacaacgaCTCTGCACC-3'
hgd35	5'-GGCGGCACAggctagctacaacgaCTGGCTCCC-3'
hgd36	5'-CGGGCGGCaggctagctacaacgaACCTGGCTC-3'
hgd37	5'-AGGGATCCAaggctagctacaacgaGAAGCAGAG-3'
hgd38	5'-GGGTAGGGAggctagctacaacgaCCATGAAGC-3'
hgd39	5'-GGGCTGAGAggctagctacaacgaTCCAGGGGG-3'
hgd40	5'-GTGGATGGAggctagctacaacgaGTCTTGAG-3'
hgd41	5'-CGTGGTGGAggctagctacaacgaGGACGTCTT-3'
hgd42	5'-GGGGGTAGAggctagctacaacgaGGAGAGGGG-3'
hgd43	5'-GGAGGAGGAggctagctacaacgaGAGGCCGGG-3'
hgd44	5'-GCCCCCGAggctagctacaacgaAAGGAGGAG-3'
hgd45	5'-CCGGGGAGAggctagctacaacgaGTCCCTTCGG-3'
hgd46	5'-GGACAGCGAggctagctacaacgaGGGTCCGGG-3'
hgd47	5'-TGGGGTGGAggctagctacaacgaAGCGATGGG-3'
hgd48	5'-CTTGAGGCaggctagctacaacgaTCTTTCTCG-3'
hgd49	5'-CACCTGGTAggctagctacaacgaTTGAGGCAC-3'

Name	DNAzyme Sequenz
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hgd51	5' -CCAGCTTCaggctagctacaacgaGCTGTCGGG-3'
hgd52	5' -GTGGGACGaggctagctacaacgaTCCAGCTTC-3'
hgd53	5' -GGAGTGGGaggctagctacaacgaGACTCCAGC-3'
hgd54	5' -ATGCTGCCAggctagctacaacgaGGGAGTGGG-3'
hgd55	5' -GGGCGGTCaggctagctacaacgaGCTGCCACG-3'
hgd56	5' -GAGGCTCCAggctagctacaacgaCCAGGGCGG-3'
hgd57	5' -GTGGGTTCGaggctagctacaacgaGAGGAGGCT-3'
hgd58	5' -AGGTGGTGAggctagctacaacgaGGGGTGGTG-3'
hgd59	5' -ACTCGGGCaggctagctacaacgaGTAGGGCGG-3'
hgd60	5' -GGAGCTGTAggctagctacaacgaTCGGGCACG-3'
hgd61	5' -GGACTTGCaggctagctacaacgaCCGAAGCCG-3'
hgd62	5' -GGGCCTGGAggctagctacaacgaTTGCATCCG-3'
hgd63	5' -TGTGCTGGAggctagctacaacgaCGGGCCTTG-3'
hgd64	5' -GTTCACACAggctagctacaacgaTCCCTGCCT-3'
hgd65	5' -CAGTTCACAggctagctacaacgaACTCCCTGC-3'
hgd66	5' -CACAGTTCaggctagctacaacgaACACTCCCT-3'
hgd67	5' -GTTGCCCCAggctagctacaacgaAGTTCACAC-3'
hgd68	5' -TCGCCGCCAggctagctacaacgaAGTGGGGTC-3'
hgd69	5' -CCCGTGCCAggctagctacaacgaCTCGCCGCC-3'
hgd70	5' -GGCGTTGCaggctagctacaacgaAGGTAGTGT-3'

### Multiple Sequence Alignments GATA-3

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Sequenz_2	****	-----	****
Sequenz_3	1	GGCGCCGCTCTTGATAC TTT CAGAAAGAATGCATTCCCTGTAAAAA	60
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Sequenz_2	****	-----	****
Sequenz_3	61	TGAGAGAGAGAGAGAGAGAGAGAGAGACGAGGGAGAGCGAGACAGAGCG	120
Sequenz_1	120	AGCAACGCAATCTGAC CGAGCAGGTCGTACGCCGCCCTCCTCTCTGCTCTTC	179
Sequenz_2	****	-----	****
Sequenz_3	121	AGCAACGCAATCTGAC CGAGCAGGTCGTACGCCGCCCTCCTCTCTCTGCTCTTC	180
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Sequenz_2	****	-----	****
Sequenz_3	181	GCTACCCAGGTGACCC GAGGAGGGACTCCGCCTCCGAGCGGCTGAGGACCCGGTGCAGA	240
Sequenz_1	240	GGAGCCTGGCTCGCAG AATTGCAGAGTCGTCGCCCTTTTACAACTGGTCCCGTTT	299
Sequenz_2	****	-----	****
Sequenz_3	241	GGAGCCTGGCTCGCAG AATTGCAGAGTCGTCGCCCTTTTACAACTGGTCCCGTTT	300
Sequenz_1	300	TTCTGCC TACCCAGT TTTTGGATTTTGTCTTCCCTTCTTCTCTTTGCTAAACGACCC	359
Sequenz_2	****	-----	****
Sequenz_3	301	TTCTGCC TACCCAGT TTTTGGATTTTGTCTTCCCTTCTTCTCTTTGCTAAACGACCC	360
Sequenz_1	360	CTCCAAGATAATTTTT AAAAACCTTCTCCTTTGCTCACCTTTGCTTCCAGCCTTCCCA	419
Sequenz_2	1	-----TCCAGCCTTCCCA	14
Sequenz_3	361	CTCCAAGATAATTTTT AAAAACCTTCTCCTTTGCTCACCTTTGCTTCCAGCCTTCCCA	420
Sequenz_1	420	TCCCCCACCAGAAAGC AAATCATTCACGACCCCGACCCCTCCGACGGCAGGAGCCCCC	479
Sequenz_2	15	TCCCCCACCAGAAAGC AAATCATTCACGACCCCGACCCCTCCGACGGCAGGAGCCCCC	74
Sequenz_3	421	TCCCCCACCAGAAAGC AAATCATTCACGACCCCGACCCCTCCGACGGCAGGAGCCCCC	480
Sequenz_1	480	GACCTCCAGCGGAC CGCCCT CCTC CGCGCGGGTTCGGGCCCGGCGAGAGGGC	539
Sequenz_2	75	GACCTCCAGCGGAC CGCCCT CC CGCGCGGGTTCGGGCCCGGCGAGAGGGC	133
Sequenz_3	481	GACCTCCAGCGGAC CGCCCT CCTC CGCGCGGGTTCGGGCCCGGCGAGAGGGC	540
Sequenz_1	540	GCGA ACAGCCGAGG CCATGGAGGTGACGGCGGACCAGCCGCGCTGGGTGAGCCACCAC	599
Sequenz_2	134	GCGA ACAGCCGAGG CCATGGAGGTGACGGCGGACCAGCCGCGCTGGGTGAGCCACCAC	193
Sequenz_3	541	GCGA ACAGCCGAGG CCATGGAGGTGACGGCGGACCAGCCGCGCTGGGTGAGCCACCAC	600
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Sequenz_2	194	CACCCCGCCGTGCTCA ACGGGCAGCACCAGGACACGCACCACCCGGGCTCAGCCACTCC	253
Sequenz_3	601	CACCCCGCCGTGCTCA ACGGGCAGCACCAGGACACGCACCACCCGGGCTCAGCCACTCC	660
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Sequenz_3	661	TACATGGACGCGGCGC AGTACCCGCTGCCGAGGAGGTGGATGTGCTTTTAAACATCGAC	720
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Sequenz_2	314	GGTCAAGGCAACCACG TCCCGCCCTACTACGAAACTCGGTACAGGGCCACGGTGCAGAGG	373
Sequenz_3	721	GGTCAAGGCAACCACG TCCCGCCCTACTACGAAACTCGGTACAGGGCCACGGTGCAGAGG	780
Sequenz_1	780	TACCTCCGACCCACC ACGGGAGCCAGGTGTGCCGCCCGCTCTGCTTCATGGATCCCTA	839
Sequenz_2	374	TACCTCCGACCCACC ACGGGAGCCAGGTGTGCCGCCCGCTCTGCTTCATGGATCCCTA	433
Sequenz_3	781	TACCTCCGACCCACC ACGGGAGCCAGGTGTGCCGCCCGCTCTGCTTCATGGATCCCTA	840
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Sequenz_2	434	CCCTGGCTGGACGGCG GCAAAGCCCTGGGACGCCACCACACCGCTCCCCCTGGAATCTC	493
Sequenz_3	841	CCCTGGCTGGACGGCG GCAAAGCCCTGGGACGCCACCACACCGCTCCCCCTGGAATCTC	900
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Sequenz_2	494	AGCCCTTCTCCAAGA CGTCCATCCACCAGGCTCCCCGGGGCCCCCTCTCCGTCTACCCC	553
Sequenz_3	901	AGCCCTTCTCCAAGA CGTCCATCCACCAGGCTCCCCGGGGCCCCCTCTCCGTCTACCCC	960
Sequenz_1	960	CCGGCCTCGTCTCTCT CTTGTGCGGGGGGCCACGCCAGCCCGACCTTTACCTTCCCG	1019
Sequenz_2	554	CCGGCCTCGTCTCTCT CTTGTGCGGGGGGCCACGCCAGCCCGACCTTTACCTTCCCG	613
Sequenz_3	961	CCGGCCTCGTCTCTCT CTTGTGCGGGGGGCCACGCCAGCCCGACCTTTACCTTCCCG	1020
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Sequenz_2	614	CCACCCCGCGGAAGG ACGTCTCCCCGGACCCATCGCTGTCCACCCAGGCTCGGCCGGC	673
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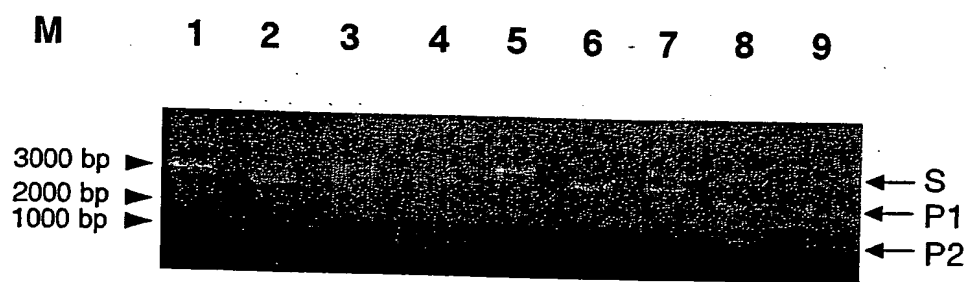
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Sequenz_3	1081	TCGGCCCCGGCAGGACG AGAAAGAGTGCCTCAAGTACCAGGTGCCCTGCCCGACAGCATG	1140
Sequenz_1	1140	AAGCTGGAGTCGTCCC ACTCCCGTGGCAGCATGACCGCCCTGGGTGGAGCCTCCTCGTCG	1199
Sequenz_2	734	AAGCTGGAGTCGTCCC ACTCCCGTGGCAGCATGACCGCCCTGGGTGGAGCCTCCTCGTCG	793
Sequenz_3	1141	AAGCTGGAGTCGTCCC ACTCCCGTGGCAGCATGACCGCCCTGGGTGGAGCCTCCTCGTCG	1200
Sequenz_1	1200	ACCCACCACCCCATCA CCACCTACCCGCCCTACGTGCCCGAGTACAGCTCCGGACTCTTC	1259
Sequenz_2	794	ACCCACCACCCCATCA CCACCTACCCGCCCTACGTGCCCGAGTACAGCTCCGGACTCTTC	853
Sequenz_3	1201	ACCCACCACCCCATCA CCACCTACCCGCCCTACGTGCCCGAGTACAGCTCCGGACTCTTC	1260
Sequenz_1	1260	CCCCCAGCAGCCTGC TGGGCGGCTCCCCACCGGCTTCGGATGCAAGTCCAGGCCCAAG	1319
Sequenz_2	854	CCCCCAGCAGCCTGC TGGGCGGCTCCCCACCGGCTTCGGATGCAAGTCCAGGCCCAAG	913
Sequenz_3	1261	CCCCCAGCAGCCTGC TGGGCGGCTCCCCACCGGCTTCGGATGCAAGTCCAGGCCCAAG	1320
Sequenz_1	1320	GCCCGGTCCAGCACAG AAGGCAGGGAGTGTGTGAACCTGTGGGGCAACCTCGACCCCACTG	1379
Sequenz_2	914	GCCCGGTCCAGCACAG ---GCAGGGAGTGTGTGAACCTGTGGGGCAACCTCGACCCCACTG	970
Sequenz_3	1321	GCCCGGTCCAGCACAG AAGGCAGGGAGTGTGTGAACCTGTGGGGCAACCTCGACCCCACTG	1380
Sequenz_1	1380	TGGCGGCGAGATGGCA CGGGACACTACCTGTGCAACGCCTGCGGGCTCTATCAGAAAATG	1439
Sequenz_2	971	TGGCGGCGAGATGGCA CGGGACACTACCTGTGCAACGCCTGCGGGCTCTATCAGAAAATG	1030
Sequenz_3	1381	TGGCGGCGAGATGGCA CGGGACACTACCTGTGCAACGCCTGCGGGCTCTATCAGAAAATG	1440
Sequenz_1	1440	AACGGACAGAACCAGC CCCTCATTAAAGCCCAAGCGAAGGCTGTCTGCAGCCAGGAGAGCA	1499
Sequenz_2	1031	AACGGACAGAACCAGC CCCTCATTAAAGCCCAAGCGAAGGCTGTCTGCAGCCAGGAGAGCA	1090
Sequenz_3	1441	AACGGACAGAACCAGC CCCTCATTAAAGCCCAAGCGAAGGCTGTCTGCAGCCAGGAGAGCA	1500
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Sequenz_2	1091	GGGACGTCTGTGCGA ACTGTGAGACCACCA CAACCACACTCTGGAGGAGGAATGCCAAT	1150
Sequenz_3	1501	GGGACGTCTGTGCGA ACTGTGAGACCACCA CAACCACACTCTGGAGGAGGAATGCCAAT	1560
Sequenz_1	1560	GGGGACCCTGTCTGCA ATGCTGTGGGCTCTACTACAAGCTTCACAATATTTAAGAGCCC	1619
Sequenz_2	1151	GGGGACCCTGTCTGCA ATGCTGTGGGCTCTACTACAAGCTTCACAATATTTAAGAGCCC	1210
Sequenz_3	1561	GGGGACCCTGTCTGCA ATGCTGTGGGCTCTACTACAAGCTTCACAATATTTAAGAGCCC	1620
Sequenz_1	1620	CTGACTATGAAGAAGG AAGGCATCCAGACCAGAAACCGAAAAATGTCTAGCAAATCCAAA	1679
Sequenz_2	1211	CTGACTATGAAGAAGG AAGGCATCCAGACCAGAAACCGAAAAATGTCTAGCAAATCCAAA	1270
Sequenz_3	1621	CTGACTATGAAGAAGG AAGGCATCCAGACCAGAAACCGAAAAATGTCTAGCAAATCCAAA	1680
Sequenz_1	1680	AAGTGCAAAAAAGTGC ATGACTCACTGGAGGACTTCCCCAAGAACAGCTCGTTTAAACCCG	1739
Sequenz_2	1271	AAGTGCAAAAAAGTGC ATGACTCACTGGAGGACTTCCCCAAGAACAGCTCGTTTAAACCCG	1330
Sequenz_3	1681	AAGTGCAAAAAAGTGC ATGACTCACTGGAGGACTTCCCCAAGAACAGCTCGTTTAAACCCG	1740
Sequenz_1	1740	GCCGCCCTCTCCAGAC ACATGTCTCCCTGAGCCACATCTCGCCCTTCAGCCACCCAGC	1799
Sequenz_2	1331	GCCGCCCTCTCCAGAC ACATGTCTCCCTGAGCCACATCTCGCCCTTCAGCCACCCAGC	1390
Sequenz_3	1741	GCCGCCCTCTCCAGAC ACATGTCTCCCTGAGCCACATCTCGCCCTTCAGCCACCCAGC	1800
Sequenz_1	1800	CACATGCTGACCACGC CCACGCCGATGCACCCGCCATCCAGCCTGTCTTTGGACCACAC	1859
Sequenz_2	1391	CACATGCTGACCACGC CCACGCCGATGCACCCGCCATCCAGCCTGTCTTTGGACCACAC	1450
Sequenz_3	1801	CACATGCTGACCACGC CCACGCCGATGCACCCGCCATCCAGCCTGTCTTTGGACCACAC	1860
Sequenz_1	1860	CACCCCTCCAGCATGG TCACCGCCATGGGTAGAGCCCTGCTCGATGCTCACAGGGCCCC	1919
Sequenz_2	1451	CACCCCTCCAGCATGG TCACCGCCATGGGTAGAGCCCTGCTCGATGCTCACAGGGCCCC	1510
Sequenz_3	1861	CACCCCTCCAGCATGG TCACCGCCATGGGTAGAGCCCTGCTCGATGCTCACAGGGCCCC	1920
Sequenz_1	1920	CAGCGAGAGTCCCTGC AGTCCCTTTTCGACTTGCATTTTTGCAGGAGCAGTATCATGAAGC	1979
Sequenz_2	1511	CAGCGAGAGTCCCTGC AGTCCCTTTTCGACTTGCATTTTTGCAGGAGCAGTATCATGAAGC	1570
Sequenz_3	1921	CAGCGAGAGTCCCTGC AGTCCCTTTTCGACTTGCATTTTTGCAGGAGCAGTATCATGAAGC	1980
Sequenz_1	1980	CTAAACGCGATGGATA TATGTTTTTGAAGGCAGAAAGCAAAATTATGTTTGCCACTTTGC	2039
Sequenz_2	1571	CTAAACGCGATGGATA TATGTTTTTGAAGGCAGAAAGCAAAATTATGTTTGCCACTTTGC	1630
Sequenz_3	1981	CTAAACGCGATGGATA TATGTTTTTGAAGGCAGAAAGCAAAATTATGTTTGCCACTTTGC	2040
Sequenz_1	2040	AAAGGAGCTCACTGTG GTGTCTGTGTTCCAACCACTGAATCTGGACCCCATCTGTGAATA	2099
Sequenz_2	1631	AAAGGAGCTCACTGTG GTGTCTGTGTTCCAACCACTGAATCTGGACCCCATCTGTGAATA	1690
Sequenz_3	2041	AAAGGAGCTCACTGTG GTGTCTGTGTTCCAACCACTGAATCTGGACCCCATCTGTGAATA	2100

Sequenz_1	2100	AGCCATTCTGACTCAT ATCCCCTATTTAACAGGGTCTCTAGTGCTGTGAAAAAAAAA-T	2158
Sequenz_2	1691	AGCCATTCTGACTCAT ATCCCCTATTTAACAGGGTCTCTAGTGCTGTGAAAAAAAAAAT	1750
Sequenz_3	2101	AGCCATTCTGACTCAT ATCCCCTATTTAACAGGGTCTCTAGTGCTGTGAAAAAAAAAAT	2160
Sequenz_1	2159	CTGAACATTGCATAT AACTTATATTGTAAGAAATACTGTACAATGACTTTATTGCATCT	2218
Sequenz_2	1751	CTGAACATTGCATAT AACTTATATTGTAAGAAATACTGTACAATGACTTTATTGCATCT	1810
Sequenz_3	2161	CTGAACATTGCATAT AACTTATATTGTAAGAAATACTGTACAATGACTTTATTGCATCT	2220
Sequenz_1	2219	GGGTAGCTGTAAGGCA TGAAGGATGCCAAGAAGTTTAAGGAATATGGGAGAAATAGTGTG	2278
Sequenz_2	1811	GGGTAGCTGTAAGGCA TGAAGGATGCCAAGAAGTTTAAGGAATATGGGAGAAATAGTGTG	1870
Sequenz_3	2221	GGGTAGCTGTAAGGCA TGAAGGATGCCAAGAAGTTTAAGGAATATGGGAGAAATAGTGTG	2280
Sequenz_1	2279	GAAATTAAGAAGAAAC TAGGTCTGATATTCAAATGGACAACTGCCAGTTTGTTCCTTT	2338
Sequenz_2	1871	GAAATTAAGAAGAAAC TAGGTCTGATATTCAAATGGACAACTGCCAGTTTGTTCCTTT	1930
Sequenz_3	2281	GAAATTAAGAAGAAAC TAGGTCTGATATTCAAATGGACAACTGCCAGTTTGTTCCTTT	2340
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Sequenz_2	1931	TCACTGGCCACAGTTG TTTGATGCATTAAAAGAAAAAATAAAAAAGAAAAAGAGAAAAG	1990
Sequenz_3	2341	TCACTGGCCACAGTTG TTTGATGCATTAAAAGAAAAAATAAAAAAGAAAAA-GAGAAAAG	2399
Sequenz_1	2399	A-----	2399
Sequenz_2	1991	AAAAAAAAAGAAAAA GTTGTAGGCGAATCATTTGTTCAAAGCTGTTGGCCCTCTGCAA	2050
Sequenz_3	2400	AAAAAAAAAGAAAAA GTTGTAGGCGAATCATTTGTTCAAAGCTGTTGGCC-TCTGCAA	2458
Sequenz_1	****	-----	****
Sequenz_2	2051	GGAAATACCAGTTCTG GGCAATCAGTGTTACCGTTCACCAGTTGCCATTGAGGGTTTCAG	2110
Sequenz_3	2459	GGAAATACCAGTTCTG GGCAATCAGTGTTACCGTTCACCAGTTGCCATTGAGGGTTTCAG	2518
Sequenz_1	****	-----	****
Sequenz_2	2111	AGAGCCTTTTCTAGG CCTACATGCTTTGTGAACAAGTCCCTGTAATTGTTGTTGTATG	2170
Sequenz_3	2519	AGAGCCTTTTCTAGG CCTACATGCTTTGTGAACAAGTCCCTGTAATTGTTGTTGTATG	2578
Sequenz_1	****	-----	****
Sequenz_2	2171	TATAATTCAAAGCACC AAAATAAGAAAAGATGTAGATTATTTTCATCATATTATACAGAC	2230
Sequenz_3	2579	TATAATTCAAAGCACC AAAATAAGAAAAGATGTAGATTATTTTCATCATATTATACAGAC	2638
Sequenz_1	****	-----	****
Sequenz_2	2231	CGAACTGTTGTATAAA TTTATTACTGCTAGTCTTAAGAACTGCTTTCTTTTCGTTTGT	2290
Sequenz_3	2639	CGAACTGTTGTATAAA TTTATTACTGCTAGTCTTAAGAACTGCTTTCTTTTCGTTTGT	2698
Sequenz_1	****	-----	****
Sequenz_2	2291	GTTTCAATATTTTCCT TCTCTCTCAATTTTCGGTTGAATAAACTAGATTACATTCAGTTG	2350
Sequenz_3	2699	GTTTCAATATTTTCCT TCTCTCTCAATTTTCGG-----	2731
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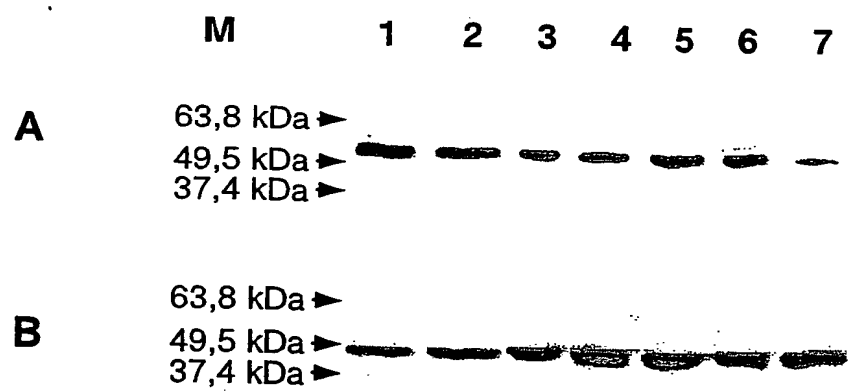
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TAAATTAATTTACTGCTACTTAAAGAACTGCTTCTTCTTCTTCT  
TTCAATTTTCTTCTCTCTCAATTTTC

Fig. 4 A





**Fig. 5**



**Fig. 6**

**Fig. 7**

Name	DNAzyme Sequenz
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td2	GGGCTCTGAggctagctacaacgaGCCTGGCTT
td3	GGGACCCCAggctagctacaacgaCGGAGCCCG
td4	GGTGGGGGAggctagctacaacgaCCCACCGGA
td5	GGCGGGGGAggctagctacaacgaCCGAGGGCC
td6	GGGCTGGGAggctagctacaacgaGGGCAGGGA
td7	CGTCGAGGAggctagctacaacgaCCGCCCCCTC
td8	GGGCTGGCAGgctagctacaacgaCTTCCCGTA
td9	CGATGCCCAGgctagctacaacgaCCGGGGCGG
td10	GCTCCACGAggctagctacaacgaGCCCATCCG
td11	CCGGCTCCAggctagctacaacgaGATGCCCAT
td12	TCTCCGCAAggctagctacaacgaCCGGCTCCA
td13	CCGTCAGCAGgctagctacaacgaGTCTCCGCA
td14	TCCCCGGCAGgctagctacaacgaCGGCTCGGT
td15	CCCCCGCGAggctagctacaacgaGCTCGTCCG
td16	GTAGGGAGAggctagctacaacgaCCCAGGCTG
td17	GGGCGGGCAGgctagctacaacgaCAAGGCGCC
td18	CGGGAAGGAggctagctacaacgaTCGCCCCGG
td19	TAGTCCTCAGgctagctacaacgaGCGGCCCCG
td20	TCCCCGACAggctagctacaacgaCTCCAGTCC
td21	TTTCCCCGAggctagctacaacgaACCTCCAGT
td22	TGAGCGCGAggctagctacaacgaCCTCAGTTT
td23	GGACCACAAGgctagctacaacgaAGGTGGTTG
td24	CTTGGAACAggctagctacaacgaAACAGGTGG
td25	AAACTTGGAggctagctacaacgaCACAACAGG
td26	CTGATTAAAggctagctacaacgaTTGGACCAC
td27	TGGTGCTGAggctagctacaacgaTAACTTGG
td28	TGATGATCAGgctagctacaacgaCTCTGTCTG
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td30	GCTTGGTGAggctagctacaacgaGATCATCTC
td31	ATGGGAACAggctagctacaacgaCCGCCGTCC
td32	GAATGGGAAggctagctacaacgaATCCGCCGT
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td36	GCCCCGCCAggctagctacaacgaAGTAAATGA
td37	CCACAAACAggctagctacaacgaCCTGTAGTG
td38	GTCCACAAAggctagctacaacgaATCCTGTAG
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td43	GCTCTGGTAggctagctacaacgaCGCCAGTGG
td44	CTGCACCCAggctagctacaacgaTTGCCGCTC
td45	CACACTGCAGgctagctacaacgaCCACTTGCC
td46	CTTTCCACAggctagctacaacgaTGCACCCAC
td47	GCCTTTCCAggctagctacaacgaACTGCACCC
td48	TTCTTGGCAGgctagctacaacgaGCTGCCCTC

Name	DNAzyme Sequenz
TD49	GTGGACGTAggctagctacaacgaAGGCGGTTT
TD50	CCGGGTGGAggctagctacaacgaGTACAGGCG
TD51	CCTGGCGCAggctagctacaacgaCCAGTGCGC
TD52	CAAATGAAAggctagctacaacgaTTCCTGGCG
TD53	TTTCCCAAaggctagctacaacgaGAAACTTCC
TD54	ATTGTTGGAggctagctacaacgaGCCCCCTTG
TD55	TGGGTCACAggctagctacaacgaTGTTGGACG
TD56	TCTGGGTCAggctagctacaacgaATTGTTGGA
TD57	GCACAATCAggctagctacaacgaCTGGGTCAC
TD58	GGAGCACAAggctagctacaacgaCATCTGGGT
TD59	ACTGGAGCAggctagctacaacgaAATCATCTG
TD60	ATGGAGGGAggctagctacaacgaTGGAGCACA
TD61	TGGTACTTAggctagctacaacgaGGAGGGACT
TD62	GGGCTGGTAggctagctacaacgaTTATGGAGG
TD63	TCAACGATAggctagctacaacgaGCAGCCGGG
TD64	CCTCAACGAggctagctacaacgaATGCAGCCG
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TD66	CGTCGTTCAggctagctacaacgaCTCAACGAT
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TD73	GCTGAGTAAggctagctacaacgaCTCGGCATT
TD74	TATTATCAaggctagctacaacgaTTTCAGCTG
TD75	GGGTTATTAggctagctacaacgaCAATTTTCA
TD76	AAGGGGTTAggctagctacaacgaTATCAATTT
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Fig. 8

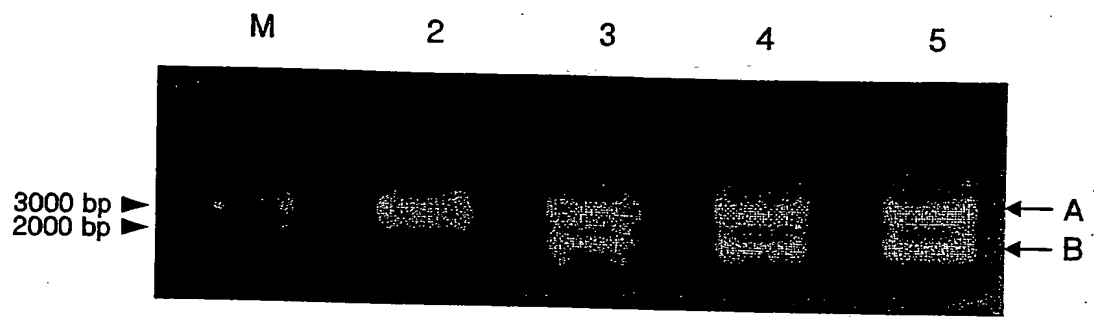
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Seq_2	61	CCAGGCGTCAGAGCCCGGGCTCCGGTGGGGTCCCCACCCGGCCCTCGGGTCCCCCGCCC	120
Seq_1	121	CCTGCTCCCTGCCATCCAGCCACGCGACCTCTCGCGCGCGGAGGGGCGGGTCTCTCG	180
Seq_2	121	CCTGCTCCCTGCCATCCAGCCACGCGACCTCTCGCGCGCGGAGGGGCGGGTCTCTCG	180
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Seq_2	181	ACGGCTACGGGAAGGTGCCAGCCCGCCCGGATGGGCATCGTGGAGCCGGGTTCGGGAGA	240
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Fig. 8A



**Fig. 9**



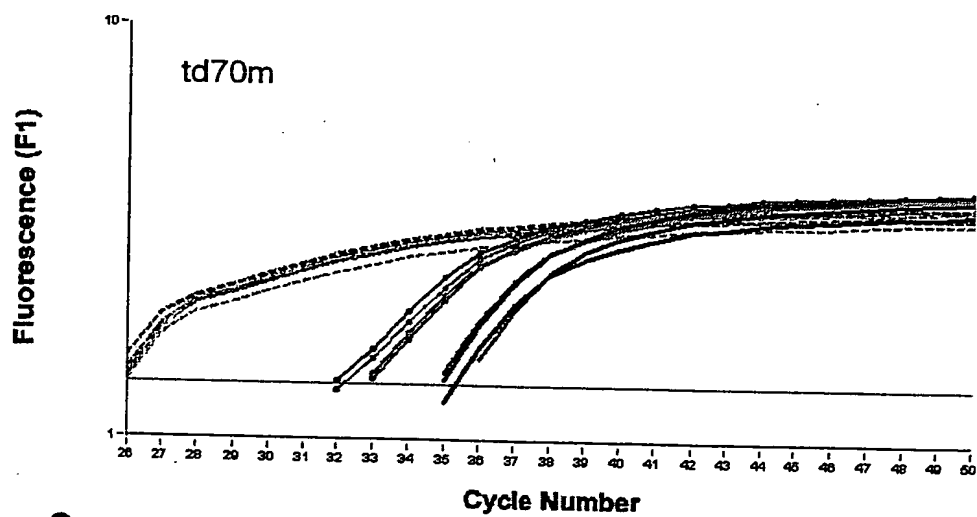
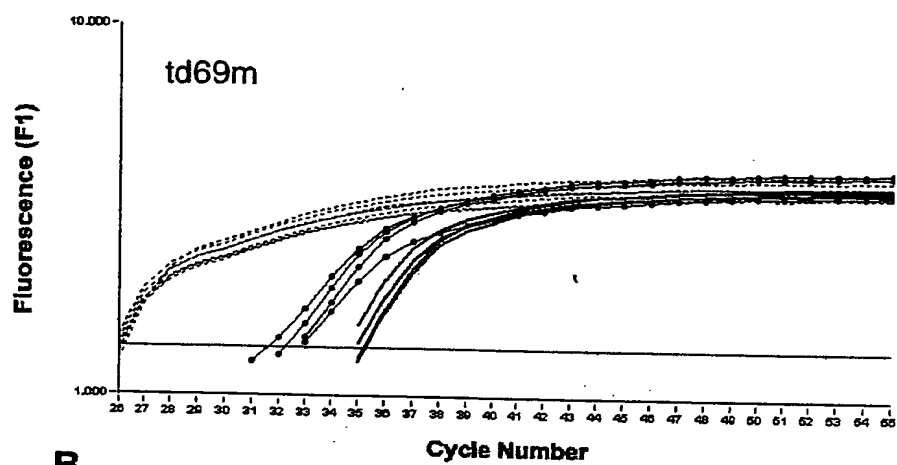
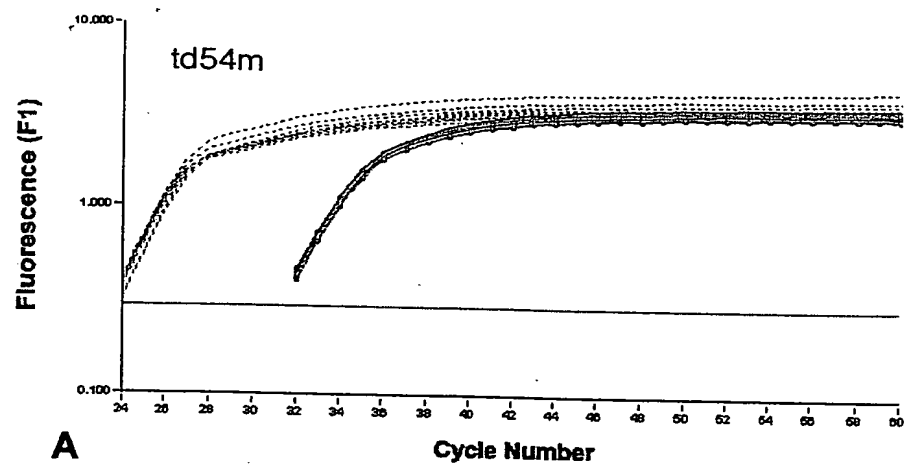
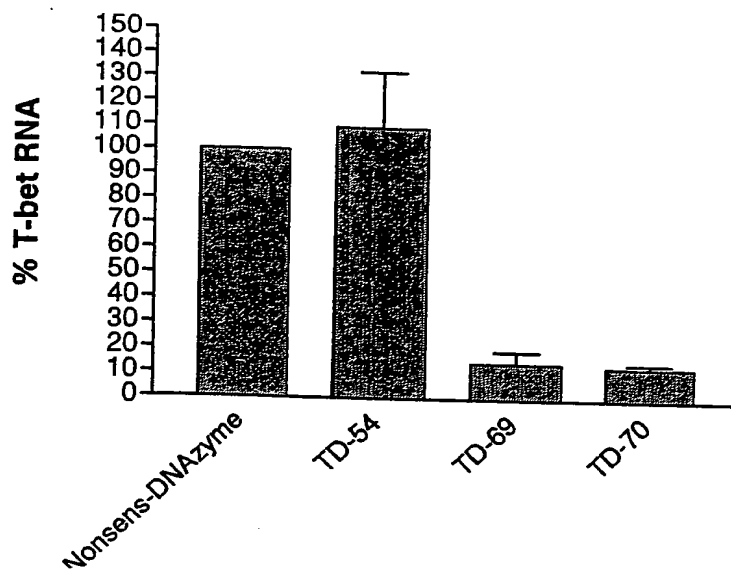


Fig. 10



**Fig. 11**